Fig 1

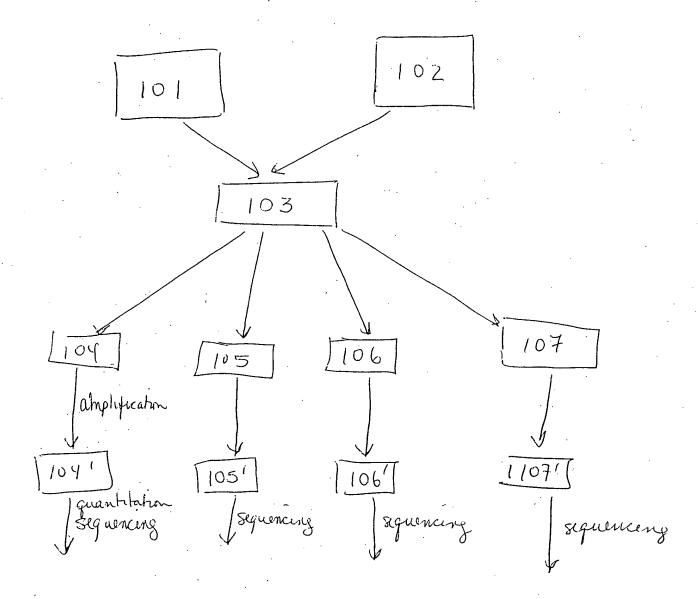
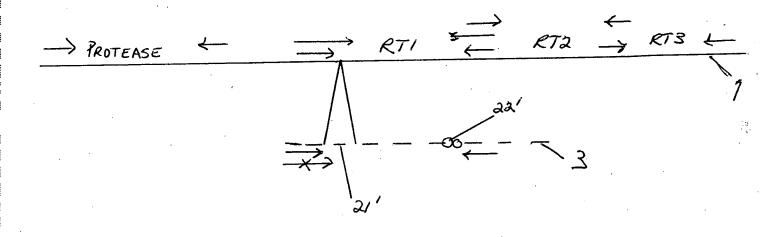


Fig 2B



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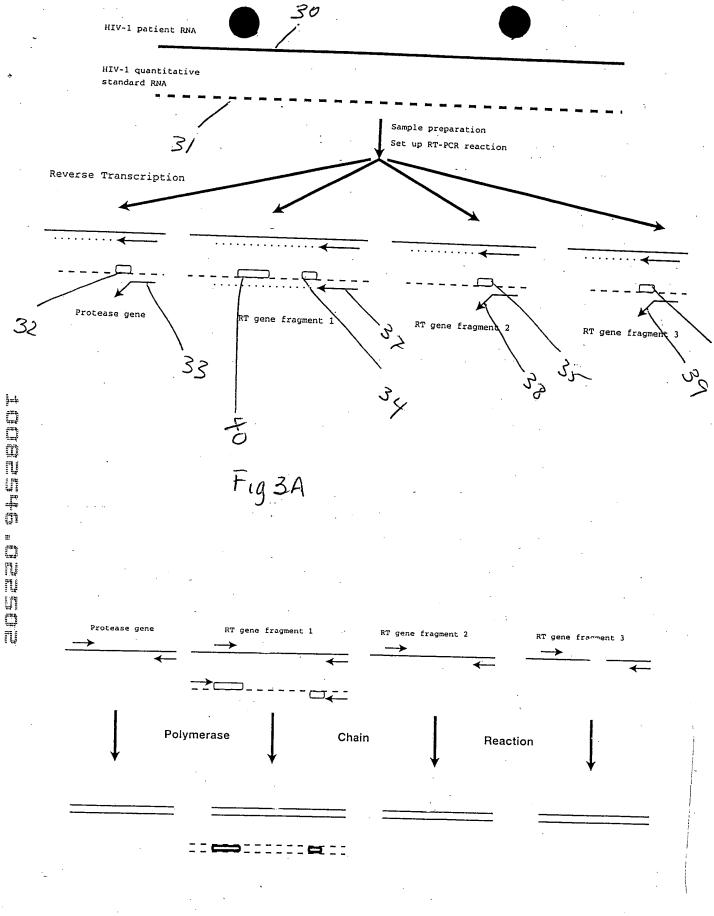
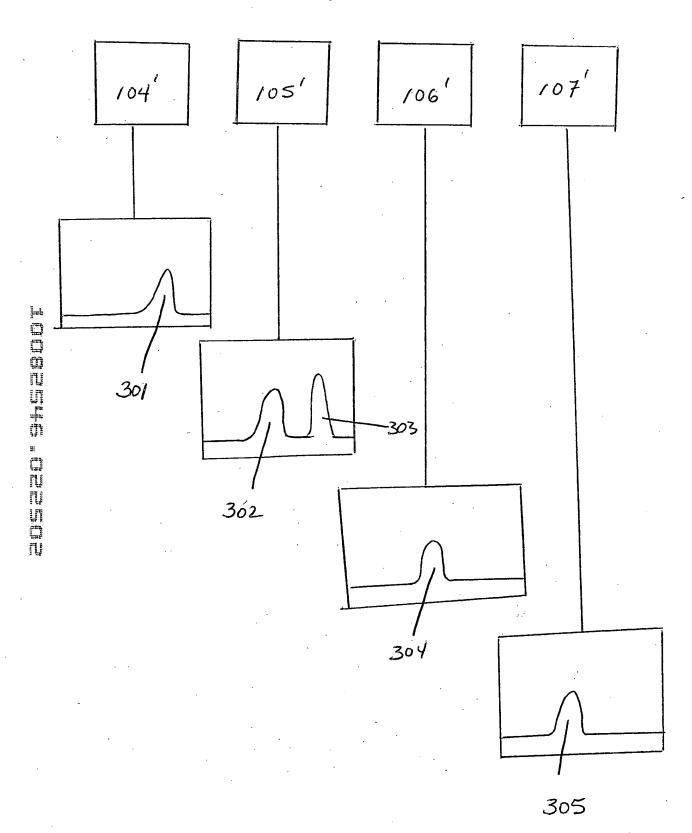


Fig 3B

Fig3C



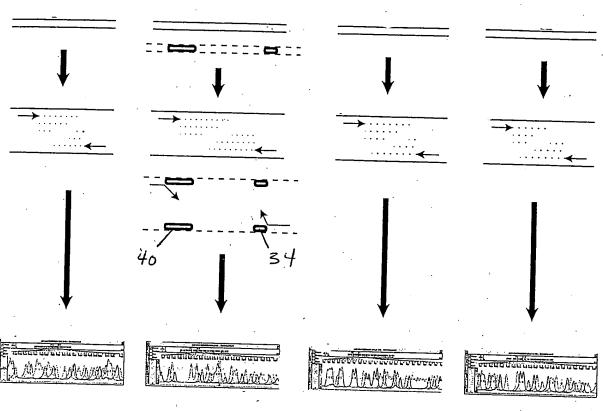
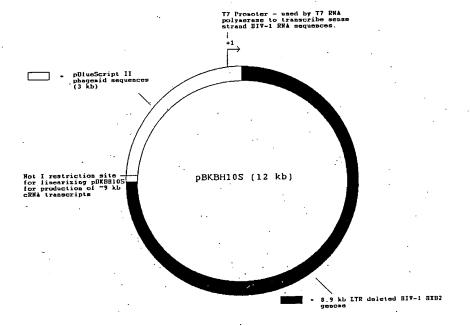


Fig 3D



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5' to 3' direction of HIV-1 sense sequence ---RT562F-A primer (for sequencing) RT558F primer (for RT-PCR) 5' aaaagttaaacaatggccattgacag aagaaaaataaaagcattagtagaaatttgtacagagatggaaaagga 3' base pairs 3' ttttcaatttgttaccggtaactgtc ttctttttattttcgtaatcatctttaaacatgtctctaccttttcct 5' 2606 to 2680 16 bp insertion added between nucleotides 2631 and 2632. The four terminal 3' nucleotides tcctaaataaatagca sequencing primer RT562F-A do not hybridize to aggatttatttatcgt sequences in the insert. Fig 5 to 3' direction of HIV-1 sense sequence _ Mutagenesis of nucleotides 3007 and 3008 CC 3007/8 gg to prevent primer extension of RT959R sequencing primer 5' tcagtacaatgtgcttccacagggat gg aaaggatcaccagcaatattccaaagtagc base pairs 3' agtcatgttacacgaaggtgtcccta cc tttcctagtggtcgttataaggtttcatcg 2981 to 3038 RT964-2-R primer (for RT-PCR) RT959R primer (for sequencing)

Figb.

	Regression Statistics	1.060750	of slope:	y-int: -0.204487	error of y-int: 0.094912	١ ١	y mean: 2.874965	x std. dev.: 0.475971	y std. dev.: 0.505588	R value: 0.998612		-		
Estimated vs. Actual HIV-1 Sample Copy Number	4		8	•		7					- 4	-	I no (Actual HIV-1 Sample	Copy Number)
					The state of the s	Log (Esumaned	HIV-1 Sample	Copy Number						

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